SEQUENCE LISTING

```
<110> Bramley, John A.
      Plaut, Karen I.
      Kerr, David
<120> TREATMENT OF STAPHYLOCOCCUS INFECTIONS
<130> Mastitis
<140> 00/000,000
<141> 2002-02-28
<160> 10
<170> PatentIn Ver. 2.1
<210> 1
<211> 1486
<212> DNA
<213> Staphylococcus simulans
<400> 1
ccggaactct tgaatgttta gttttgaaaa ttccaaaaaa aaacctacct tcttaatatt 60
gattcatatt attttaacac aatcagttag aatttcaaaa atcttaaagt caatttttga 120
gtgtgtttgt atatttcatc aaaatcaatc aatattattt tactttcttc atcgttaaaa 180
aatgtaatat ttataaaaat atgctattct cataaatgta ataataaatt aggaggtatt 240
aaggttgaag aaaacaaaaa acaattatta tacgagacct ttagctattg gactgagtac 300
atttgcctta gcatctattg tttatggagg gattcaaaat gaaacacatg cttctgaaaa 360
aagtaatatg gatgtttcaa aaaaagtagc tgaagtagag acttcaaaag ccccagtaga 420
aaatacagct qaaqtagaga cttcaaaaagc tccagtagaa aatacagctg aagtagagac 480
ttcaaaaget ccagtagaaa atacagetga agtagagaet tcaaaagete cagtagaaaa 540
tacagotgaa gtagagaott caaaagotoo ggtagaaaat acagotgaag tagagaotto 600
aaaagcccca gtagaaaata cagctgaagt agagacttca aaagccctgg ttcaaaatag 660
aacagettta agagetgeaa cacatgaaca tteageacaa tggttgaata attacaaaaa 720
aggatatggt tacggtcctt atccattagg tataaatggc ggtatgcact acggagttga 780
ttttttttatg aatattggaa caccagtaaa agctatttca agcggaaaaa tagttgaagc 840
tggttggagt aattacggag gaggtaatca aataggtctt attgaaaatg atggagtgca 900
tagacaatgg tatatgcatc taagtaaata gtaggagatt gtaggagatt atgtcaaagc 960
tggtcaaata atcggttggt ctggaagcac tggttattct acagcaccac atttacactt 1020
ccaaagaatg gttaattcat tttcaaattc aactgcccaa gatccaatgc ctttcttaaa 1080
gagcgcagga tatggaaaag caggtggtac agtaactcca acgccgaata caggttggaa 1140
aacaaacaaa tatggcacac tatataaatc agagtcagct agcttcacac ctaatacaga 1200
tataataaca agaacgactg gtccatttag aagcatgccg cagtcaggag tcttaaaagc 1260
aggtcaaaca attcattatg atgaagtgat gaaacaagac ggtcatgttt gggtaggtta 1320
tacaggtaac agtggccaac gtatttactt gcctgtaaga acatggaata aatctactaa 1380
tactttaggt gttctttggg gaactataaa gtgagcgcgc tttttataaa cttatatgat 1440
```

1486

aattagagca aataaaaatt ttttctcatt cctaaagttg aagctt

<210> 2

<211> 388

<212> PRT

<213> Staphylocccus simulans

<400> 2

Met Lys Lys Thr Lys Asn Asn Tyr Tyr Thr Arg Pro Leu Ala Ile Gly
1 5 10 15

Leu Ser Thr Phe Ala Leu Ala Ser Ile Val Tyr Gly Gly Ile Gln Asn 20 25 30

Glu Thr His Ala Ser Glu Lys Ser Asn Met Asp Val Ser Lys Lys Val
35 40 45

Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val
50 55 60

Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser
65 70 75 80

Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Pro 85 90 95

Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn 100 105 110

Thr Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu
115 120 125

Val Glu Thr Ser Lys Ala Leu Val Gln Asn Arg Thr Ala Leu Arg Ala 130 135 140

Ala Thr His Glu His Ser Gln Trp Leu Asn Asn Tyr Lys Lys Gly Tyr 145 150 155 160

Gly Tyr Gly Pro Tyr Pro Leu Gly Ile Asn Gly Gly Met His Tyr Gly
165 170 175

Val Asp Glu Phe Met Asn Ile Gly Thr Pro Val Lys Ala Ile Ser Ser 180 185 190

Gly Lys Ile Val Glu Ala Gly Trp Ser Asn Tyr Gly Gly Asn Gln 195 200 205

Ile Gly Leu Ile Glu Asn Asp Gly Val His Arg Gln Glu Tyr Met His

210 215 220

Leu Ser Lys Tyr Asn Val Lys Val Gly Asp Tyr Val Lys Ala Gly Gln 225 230 235 240

Ile Ile Gly Trp Ser Gly Ser Thr Gly Tyr Ser Thr Ala Pro His Leu 245 250 255

His Phe Gln Arg Met Val Asn Ser Phe Ser Asn Ser Thr Ala Gln Asp 260 265 270

Pro Met Pro Phe Leu Lys Ala Ser Gly Tyr Gly Lys Ala Gly Gly Thr 275 280 285

Val Thr Pro Thr Pro Asn Thr Gly Trp Lys Thr Asn Lys Tyr Gly Thr 290 295 300

Leu Tyr Lys Ser Glu Ser Ala Ser Phe Thr Pro Asn Thr Asp Ile Ile 305 310 315 320

Thr Arg Thr Thr Gly Pro Phe Arg Ser Met Pro Gln Ser Gly Val Leu 325 330 335

Lys Ala Gly Gln Thr Ile His Tyr Asp Glu Val Met Lys Gln Asp Gly 340 350

His Val Trp Val Gly Tyr Thr Gly Asn Ser Gly Gln Arg Ile Tyr Leu 355 360 365

Pro Val Arg Thr Trp Asn Lys Ser Thr Asn Thr Leu Gly Val Leu Trp 370 375 380

Gly Thr Ile Lys 385

<210> 3

<211> 741

<212> DNA

<213> Staphylococcus simulans

<400> 3

gccgcaacac atgaacattc agcacaatgg ttgaataatt acaaaaaagg atatggttac 60 ggcccttatc cattaggtat aaatggcggt atgcactacg gagttgattt ttttatgaat 120 attggaacac cagtaaaagc tatttcaagc ggaaaaatag ttgaagctgg ttggagtaat 180 tacggaggag gtaatcaaat aggtcttatt gaaaatgatg gagtgcatag acaatggtat 240 atgcatctaa gtaaatataa tgttaaagta ggagattatg tcaaagctgg tcaaataatc 300 ggttggtctg gaagcactgg ttattctaca gcaccacatt tacacttcca aagaatggtt 360

```
aactcatttt cacagtcaac tgcccaagat ccaatgcctt tcttaaagag cgcaggatat 420
ggaaaagcag gtggtacagt aactccaacg ccgaatacag gttggaaaac aaacaaatat 480
ggcacactat ataaatcaga gtcagctagc ttcacaccta atacagatat aataacaaga 540
acgactggtc catttagaag catgccgcag tcaggagtct taaaagcagg tcaaaacaatt 600
cattatgatg aagtgatgaa acaagacggt catgtttggg taggttatac aggtaacagt 660
ggccaacgta tttacttgcc tgtgagaaca tggcagaagt ctactaatac tctgggtgtt 720
                                                                  741
ctgtggggaa ctataaagtg a
<210> 4
<211> 1520
<212> DNA
<213> Staphylococcus simulans
<400> 4
tgtgtgcgtg ctcccattcg ttcatgctcg ccacgcgcac ggccgcgctt tgcgacgcga 60
tegegeaceg tgtgaacege attgaggaat ggeegttegg caagegeatg taeggeeteg 120
atttgaacgt gcgtcgcacg acagcgtcgc gcccgcggtc agagtccggc gcccgcggta 180
tacggacage gategeggeg teegeegatg acgaacggte gtgegegtea gtegeatgeg 240
ccgctcgccg ctggcgttcc ggcttcgcgg gcgcagcgcg gtccaccact cttcaaacgt 300
ctttctcggg agcagcatat gaagaagatt tccaaggcgg gactggggct ggcgctggtg 360
tgcgcgctgg cgacgatcgg cggcaacgca gcgcgcaggg ccacggctca gcggcgagga 420
tctggtgtat tctacgacga gatgttcgac ttcgacatcg atgcgcatct ggccaagcat 480
gcgccgcatc tgcacaagca ctcggaagag atctcgcact gggccggcta cagcgggatc 540
agcegaagtg ttgategege tgatggagea geagagegeg eggteaegee aagegegega 600
cgaatcgtcc gttcggcaag ctggcgcgc ccgacggctt cggcgcgcag acccgcgagg 660
tegegetgge getgegegag tegetgtaeg agegegatee egaegegeea aggggeeggt 720
gacgctggcc cgcgccaatc cgctgcaggc gctgttcgag cgttccggcg acaacgagcc 780
ggcggccgcg ctgcgcggcg acggcgagtt ccagctggtc tacggccgcc tgttcaacga 840
accgcgccag gccaaggcgg cttcggaccg cttcgccaag gccggcccgg acgtgcagcc 900
gtgtcgccca acggcctgct gcagttcccc ttcccgcgcg gcgccagctg gcatgtcggc 960
ggcgcccaca ccaacacegg ctcgggcaat tacccgatgt cgtcgctgga catgtcgcgc 1020
ggcggcggct ggggcagcaa ccagaacggc aactgggtgt cggcctcggc cgccggctcg 1080
ttcaagcgcc actcttcgtg cttcgcggag atcgtgcaca ccggcggctg gtcgacgacc 1140
tactaccacc tgatgaacat ccagtacaac accggcgcca acgtgtcgat gaacaccgcc 1200
ategecaace eggecaacae eeaggegeag gegetgtgea aeggeggeea gtegaeegge 1260
ccgcacgagc attggtcgtt gaagcagaac ggcagcttct accacctcaa cggcacctac 1320
ctgtcgggct atcgcatcac cgcgaccggc agcagctatg acaccaactg cagccggttc 1380
tatctgacca agaacggcca gaactactgc tacggctatt acgtcaaccc gggcccgaac 1440
tgaggetege egegtgegtt geeegegtee teaagegeee caegegeggg gegegggeae 1500
```

cggccgggtc aggtcgaatt

<210> 5

<211> 480

<212> PRT

<213> Staphylococcus simulans

<400 Met		Lys	Thr	Lys 5	Asn	Asn	Tyr	Tyr	Thr	Thr	Pro	Leu	Ala	Ile 15	Gly
	Ser	Thr	Phe 20	Ala	Leu	Ala	Ser	Ile 25	Val	Tyr	Gly	Gly	Ile 30	Gln	Asn
Glu	Thr	His		Ser	Glu	Lys	Ser		Met	Asp	Val	Ser		Lys	Val
Ala	Glu 50		Glu	Thr	Ser	Lys 55		Pro	Val	Glu	Asn 60		Ala	Glu	Val
Glu 65		Ser	Lys	Ala	Pro		Glu	Asn	Thr	Ala 75		Val	Glu	Thr	Ser 80
	Ala	Pro	Val	Glu 85		Thr	Ala	Glu	Val 90		Thr	Ser	Lys	Ala 95	
Val	Glu	Asn	Thr		Glu	Val	Glu	Thr		Lys	Ala	Pro	Val	Glu	Asn
Thr	Ala	Glu 115		Glu	Thr	Ser	Lys 120		Pro	Val	Glu	Asn 125		Ala	Glu
Val	Glu 130		Ser	Lys	Ala	Pro 135		Glu	Asn	Thr	Ala 140		Val	Glu	Thr
		Ala	Pro	Val			Thr	Ala	Glu			Thr	Ser	Lys	Ala 160
145 Pro	Val	Glu	Asn		150 Ala	Glu	Val			155 Ser	Lys	Ala	Pro	Val	
Asn	Thr	Ala		165 Val	Glu	Thr	Ser	Lys	170 Ala	Pro	Val	Glu		175 Thr	Ala
Glu	Val		180 Thr	Ser	Lys	Ala		185 Val	Glu	Asn	Thr		190 Glu	Val	Glu
Thr		195 Lys	Ala	Pro	Val		200 Asn	Thr	Ala	Glu		205 Glu	Thr	Ser	Lys
	210 Leu	Val	Gln	Asn		215 Thr	Ala	Leu	Arg		220 Ala	Thr	His	Glu	
225 Ser	Ala	Gln	Trp	Leu	230 Asn	Asn	Tyr	Lys	Tyr	235 Gly	Tyr	Gly	Tyr	Gly	240 Pro

Tyr	Pro	Leu	Gly 260	Ile	Asn	Gly	Gly	Ile 265	His	Tyr	Gly	Val	Asp 270	Phe	Phe
Met	Asn	Ile	Gly	Thr	Pro	Val	Lys	Ala	Ile	Ser	Ser	Gly	Lys	Ile	Val

Glu Ala Gly Trp Ser Asn Tyr Gly Gly Gly Asn Gln Ile Gly Leu Ile 290 295 300

285

280

- Glu Asn Asp Gly Val His Arg Gln Trp Tyr Met His Leu Ser Lys Tyr 305 310 315 320
- Asn Val Lys Val Gly Asp Tyr Val Lys Ala Gly Gln Ile Ile Gly Trp 325 330 335
- Ser Gly Ser Thr Gly Tyr Ser Thr Ala Pro His Leu His Phe Gln Arg 340 345 350
- Met Val Asn Ser Phe Ser Asn Ser Thr Ala Gln Asp Pro Met Pro Phe 355 360 365
- Leu Lys Ser Ala Gly Tyr Gly Lys Ala Gly Gly Thr Val Thr Pro Thr 370 375 380
- Pro Asn Thr Gly Trp Lys Thr Asn Lys Tyr Gly Thr Leu Tyr Lys Ser 385 390 395 400
- Glu Ser Ala Ser Phe Thr Pro Asn Thr Asp Ile Ile Thr Arg Thr Thr 405 410 415
- Gly Pro Phe Arg Ser Met Pro Gln Ser Gly Val Leu Lys Ala Gly Gln
 420 425 430
- Thr Ile His Tyr Asp Glu Val Met Lys Gln Asp Gly His Val Trp Val 435 440 445
- Gly Tyr Thr Gly Asn Ser Gly Gln Arg Ile Tyr Leu Pro Val Arg Thr 450 455 460
- Trp Asn Lys Ser Thr Asn Thr Leu Gly Val Leu Trp Gly Thr Ile Lys 465 470 475 480

<211> 492 <212> PRT

<213> Achromobacter lyticus

<400> б

Met Lys Lys Thr Lys Asn Asn Tyr Tyr Thr Arg Pro Leu Ala Ile Gly
1 5 10 15

Leu Ser Thr Phe Ala Leu Ala Ser Ile Val Tyr Gly Gly Ile Gln Asn 20 25 30

Glu Thr His Ala Ser Glu Lys Ser Asn Met Asp Val Ser Lys Lys Val
35 40 45

Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val
50 55 60

Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser
65 70 75 80

Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Pro
85 90 95

Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn 100 105 110

Thr Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu 115 120 125

Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr 130 135 140

Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala 145 150 155 160

Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu 165 170 175

Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala 180 185 190

Glu Val Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu
195 200 205

Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys 210 215 220

Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser Lys Ala Leu Val

Gln Arg Thr Ala Leu Arg Ala Ala Thr His Glu His Ser Ala Gln Trp
245 250 255

Leu Asn Asn Tyr Lys Lys Gly Tyr Gly Tyr Gly Pro Tyr Pro Leu Gly
260 265 270

Ile Asn Gly Gly Met His Tyr Gly Val Asp Phe Phe Met Asn Ile Gly 275 280 285

Thr Pro Val Lys Ala Ile Ser Ser Gly Lys Ile Val Glu Ala Gly Trp 290 295 300

Ser Asn Tyr Gly Gly Gly Asn Gln Ile Gly Leu Ile Glu Asn Asp Gly 305 310 315 320

Val His Arg Gln Trp Tyr Met His Leu Ser Lys Tyr Asn Val Lys Val
325 330 335

Gly Asp Tyr Val Lys Ala Gly Gln Ile Ile Gly Trp Ser Gly Ser Thr 340 345 350

Gly Tyr Ser Thr Ala Pro His Leu His Phe Gln Arg Met Val Asn Ser 355 360 365

Phe Ser Asn Ser Thr Ala Gln Asp Pro Met Pro Phe Leu Lys Ser Ala 370 375 380

Gly Tyr Gly Lys Ala Gly Gly Thr Val Thr Pro Thr Pro Asn Thr Gly 385 390 395 400

Trp Lys Thr Asn Lys Tyr Gly Thr Leu Tyr Lys Ser Glu Ser Ala Ser \$405\$

Phe Thr Pro Asn Thr Asp Ile Ile Thr Arg Thr Thr Gly Pro Phe Arg 420 425 430

Ser Met Pro Gln Ser Gly Val Leu Lys Ala Gly Gln Thr Ile His Tyr 435 440 445

Asp Glu Val Met Lys Gln Asp Gly His Val Trp Val Gly Tyr Thr Gly 450 455 460

Asn Ser Gly Gln Arg Ile Tyr Leu Pro Val Arg Thr Trp Asn Lys Ser 465 470 475 480

Thr Asn Thr Leu Gly Val Leu Trp Gly Thr Ile Lys

485 490

```
<210> 7
<211> 741
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: altered S.
      simulans lysostaphin gene
<400> 7
gctgcaacac atgaacattc agcacaatgg ttgaataatt acaaaaaagg atatggttac 60
gqtccttatc cattaggtat aaatggcggt atgcactacg gagttgattt ttttatgaat 120
attggaacac cagtaaaagc tatttcaagc ggaaaaatag ttgaagctgg ttggagtaat 180
tacggaggag gtaatcaaat aggtcttatt gaaaatgatg gagtgcatag acaatggtat 240
atgcatctaa gtaaatataa tgttaaagta ggagattatg tcaaagctgg tcaaataatc 300
ggttggtctg gaagcactgg ttattctaca gcaccacatt tacacttcca aagaatggtt 360
aattcatttt caaattcaac tgcccaagat ccaatgcctt tcttaaagag cgcaggatat 420
ggaaaagcag gtggtacagt aactccaacg ccgaatacag gttggaaaac aaacaaatat 480
ggcacactat ataaatcaga gtcagctagc ttcacaccta atacagatat aataacaaga 540
acgactggtc catttagaag catgccgcag tcaggagtct taaaagcagg tcaaacaatt 600
cattatgatg aagtgatgaa acaagacggt catgtttggg taggttatac aggtaacagt 660
ggccaacgta tttacttgcc tgtaagaaca tggaataaat ctactaatac tttaggtgtt 720
ctttggggaa ctataaagtg a
<210> 8
<211> 480
<212> PRT
<213> Staphylococcus simulans
<400> 8
Met Lys Lys Thr Lys Asn Asn Tyr Tyr Thr Thr Pro Leu Ala Ile Gly
  1
                  5
                                     10
                                                         15
Leu Ser Thr Phe Ala Leu Ala Ser Ile Val Tyr Gly Ile Gln Asn
             20
                                 25
Glu Thr His Ala Ser Glu Lys Ser Asn Met Asp Val Ser Lys Lys Val
         35
                             40
                                                 45
Ala Glu Val Glu Thr Ser Lys Pro Pro Val Glu Asn Thr Ala Glu Val
Glu Thr Ser Lys Ala Pro Val Glu Asn Thr Ala Glu Val Glu Thr Ser
```

741

80

75

L	ys	Ala	Pro	Val	Glu 85	Asn	Thr	Ala	Glu	Val 90	Glu	Thr	Ser	Lys	Ala 95	Pro
V	al	Glu	Asn	Thr 100	Ala	Glu	Val	Glu	Thr 105	Ser	Lys	Ala	Pro	Val 110	Glu	Asn
T	hr	Ala	Glu 115	Val	Glu	Thr	Ser	Lys 120	Ala	Pro	Val	Glu	Asn 125	Thr	Ala	Glu
V	al	Glu 130	Thr	Ser	Lys	Ala	Pro 135	Val	Glu	Asn	Thr	Ala 140	Glu	Val	Glu	Thr
	er 45	Lys	Ala	Pro	Val	Glu 150	Asn	Thr	Ala	Glu	Val 155	Glu	Thr	Ser	Lys	Ala 160
P:	ro	Val	Glu	Asn	Thr 165	Ala	Glu	Val	Glu	Thr 170	Ser	Lys	Ala	Pro	Val 175	Glu
A	sn	Thr	Ala	Glu 180	Val	Glu	Thr	Ser	Lys 185	Ala	Pro	Val	Glu	Asn 190	Thr	Ala
G.	lu	Val	Glu 195	Thr	Ser	Lys	Ala	Pro 200	Val	Glu	Asn	Thr	Ala 205	Glu	Val	Glu
T]	hr	Ser 210	Lys	Ala	Pro	Val	Glu 215	Asn	Thr	Ala	Glu	Val 220	Glu	Thr	Ser	Lys
	1a 25	Leu	Val	Gln	Asn	Arg 230	Thr	Ala	Leu	Arg	Ala 235	Ala	Thr	His	Glu	His 240
S	er	Ala	Gln	Trp	Leu 245	Asn	Asn	Tyr	Lys	Tyr 250	Gly	Tyr	Gly	Tyr	Gly 255	Pro
T	yr	Pro	Leu	Gly 260	Ile	Asn	Gly	Gly	Ile 265	His	Tyr	Gly	Val	Asp 270	Phe	Phe
Me	et	Asn	Ile 275	Gly	Thr	Pro	Val	Lys 280	Ala	Ile	Ser	Ser	Gly 285	Lys	Ile	Val
G.	lu	Ala 290	Gly	Trp	Ser	Asn	Tyr 295	Gly	Gly	Gly	Asn	Gln 300	Ile	Gly	Leu	Ile
	lu 05	Asn	Asp	Gly	Val	His 310	Arg	Gln	Trp	Tyr	Met 315	His	Leu	Ser	Lys	Tyr 320
A	sn	Val	Lys	Val	Gly	Asp	Tyr	Val	Lys	Ala	Gly	Gln	Ile	Ile	Gly	Trp

```
Ser Gly Ser Thr Gly Tyr Ser Thr Ala Pro His Leu His Phe Gln Arg
340 345 350
```

Met Val Asn Ser Phe Ser Asn Ser Thr Ala Gln Asp Pro Met Pro Phe 355 360 365

Leu Lys Ser Ala Gly Tyr Gly Lys Ala Gly Gly Thr Val Thr Pro Thr 370 375 380

Pro Asn Thr Gly Trp Lys Thr Asn Lys Tyr Gly Thr Leu Tyr Lys Ser 385 390 395 400

Glu Ser Ala Ser Phe Thr Pro Asn Thr Asp Ile Ile Thr Arg Thr Thr
405 410 415

Gly Pro Phe Arg Ser Met Pro Gln Ser Gly Val Leu Lys Ala Gly Gln
420 425 430

Thr Ile His Tyr Asp Glu Val Met Lys Gln Asp Gly His Val Trp Val 435 440 445

Gly Tyr Thr Gly Asn Ser Gly Gln Arg Ile Tyr Leu Pro Val Arg Thr 450 455 460

Trp Asn Lys Ser Thr Asn Thr Leu Gly Val Leu Trp Gly Thr Ile Lys 465 470 475 480

<210> 9

<211> 1825

<212> DNA

<213> Staphylococcus simulans

<400> 9

gaaaattcca aaaaaaaacc tacttctta atattgattc atattattt aacacaatca 60 gttagaattt caaaaatctt aaagtcaatt tttgagtgtg tttgtatatt tcatcaaagc 120 caatcaatat tattttactt tcttcatcgt taaaaaaatgt aatattata aaaatatgct 180 attctcataa atgtaataat aaattaggag gtattaaggt tgaagaaaac aaaaaacaat 240 tattatacga cacctttagc tattggactg agtacatttg ccttagcatc tattgtttat 300 ggagggattc aaaatgaaac acatgcttct gaaaaaaagta atatggatgt ttcaaaaaaaa 360 gtagctgaag tagagacttc aaaaccccca gtagaaaata cagctgaagt agaacttca 420 aaagctccag tagaaaatac agctgaagta gagacttcaa aagctccagt agaaaataca 480 gctgaagtag agacttcaaaa agctccagta gaaaatacag ctgaagtaga gacttcaaaa 540 gctccggtag aaaatacagc tgaagtagag acttcaaaag ctccggtaga aaaatacagc 600

```
qaaqtagaga cttcaaaagc cccagtagaa aatacagctg aagtagagac ttcaaaagct 660
ccagtagaaa atacagctga agtagagact tcaaaaagctc cggtagaaaa tacagctgaa 720
gtagagactt caaaagcccc agtagaaaat acagctgaag tagagacttc aaaagctcca 780
gtagaaaata cagctgaagt agagacttca aaagctccgg tagaaaatac agctgaagta 840
gagacttcaa aagccccagt agaaaataca gctgaagtag agacttcaaa agccctggtt 900
caaaatagaa cagctttaag agctgcaaca catgaacatt cagcacaatg gttgaataat 960
tacaaaaaag gatatggtta cggtccttat ccattaggta taaatggcgg tatccactac 1020
ggagttgatt tttttatgaa tattggaaca ccagtaaaag ctatttcaag cggaaaaata 1080
gttgaagctg gttggagtaa ttacggagga ggtaatcaaa taggtcttat tgaaaatgat 1140
ggagtgcata gacaatggta tatgcatcta agtaaatata atgttaaagt aggagattat 1200
gtcaaagctg gtcaaataat cggttggtct ggaagcactg gttattctac agcaccacat 1260
ttacacttcc aaagaatggt taattcattt tcaaattcaa ctgcccaaga tccaatgcct 1320
ttcttaaaga gcgcaggata tggaaaagca ggtggtacag taactccaac gcccaataca 1380
ggttggaaaa caaacaaata tggcacacta tataaatcag agtcagctag cttcacacct 1440
aatacagata taataacaag aacgactggt ccatttagaa gcatgccgca gtcaggagtc 1500
ttaaaagcag gtcaaacaat tcattatgat gaagtgatga aacaagacgg tcatgtttgg 1560
gtaggttata caggtaacag tggccaacgt atttacttgc ctgtaagaac atggaataaa 1620
tctactaata ctttaggtgt tctttgggga actataaagt gagcgcgctt tttataaact 1680
tatatgataa ttagagcaaa taaaaatttt ttctcattcc taaagttgaa gcttttcgta 1740
atcatgtcat agcgtttcct gtgtgaaatt gcttagcctc acaattccac acaacatacg 1800
                                                                  1825
agccggaaca taaagtgcta agcct
```

<210> 10 <211> 6457 <212> DNA <213> Achromobacter lyticus

<400> 10

gatatcattt caaagacaga tattctaaag aaaagatata ttttaaaaaa tgtggttgaa 60 aaaattaaag aaattcacga ttttgactat atatttattg atgtaccacc tactattaac 120 tctgatttca ctaataatgc tgtttacgca agtgattaca ttttaatggt atttcaaaca 180 caacaatctg cttatgaaag tagtctttca tttgttaatt ttttaaggga tcgaaaaaaa 240 gaatcagatt tatcatttga attggttggc gctgttccag tattaattaa aaaaagtgga 300 cgtgtagata aacagatatt agatatgtct aaatcagcat tttctgaagc actctttgag 360 aaccagatat atcaaagaga aagaataaaa aaatttgccg ctgatggaat aaaagataaa 420 gatatgcatg acaaaaaagt tatatatatg tttaacaaag tctacgaaga attagttgat 480 agagttagat taattgaagg tgagtgatat ttatggcagg atttttagat aacatagata 540 catctgaggt aaaatatacg gaaaattata aaccggtatc taaaagtacg actatgagag 600 tggacactga tataaaaaaa agattaaatc aaatggcgtt agataaagat acatctataa 660 aggctatagt tgatgaagtg ttaggagaat ttttgaaaaa aaataagtat tagtatttta 720 tataggetet ataetattta ggaetggtga taateaetag teetattttt gataeaaaaa 780 agcgcaatta tctctataat tagaagtatc ctaccaccaa taattaagga aataatgcgc 840 ctatgtctaa tattatatca atcaccettg gaattaaaga taaaaatatc acttttgaag 900 ataaggttga agaaagtata aagggaaaaa ttctttattt tactttggaa aattaataca 960 ttctcccaag cgatgtaaac tttgcggaca cgaaaatacg aacttttcta taatcaaaaa 1020 tggttttaaa aaatcatgtc ttacgatacc taaggtatcg gagaagccag cttatttaat 1080 attggaaaaa cagcgtttcc actgtaaaaa gtgctgcagt tatttcactg ctgaaacacc 1140

tgtcgttgag tggaattgct atatttctca aaacacacga ttagctgtgc tgaataagtc 1200 gatagacata cgttcgcaaa aatctgttgc tgaatcttgt catgtcagta attccacagt 1260 tactogaata attaataaag otgottotoa aatagotoaa acacogttta aatatttaco 1320 ggaacacttg atgatggatg agttcaaaag cgttaaaaat gttgtcggta aaatgagttt 1380 tatttatgca gatgcagtaa cacaccgtat tattgatatt gtgcctgacc gcaggttatt 1440 tgctttgaaa aattatttct accgttatcc tctttctgaa agaaaatgtg tgaaagcagt 1500 gtctattgat atgtatgaac cttatatggc tttgatcaga gaagtttttc ctaatgccaa 1560 aattctaata gttcatttcc atattgttca gtctttaaat aaagccttga acatgactcg 1620 agtaacagtt atgaatagtt tcagaacaac tgaaagacct ctatacaaca agtacaagcg 1680 ttactggaag attctttaa aactgccttg aaaaatatag aaatcaatag cgttgctcct 1740 aaacttcaaa cagctgttaa aacactaaga aagcacaata gaatgataag aaatactttt 1800 gaatacagta acttgaccaa cggttcactt gagggaataa atactaaaat aaagctgata 1860 cagagaatat cttttggtta tagaaatttt ggtgatttac gcagtcgtat cattttatgt 1920 acaaatcttt ttgcagctaa tccaaaaaaa gagatcaagc aactttatgc tgcttaatct 1980 ctgcgtttta gctcaccagt cttatttgac agagagccaa taaaattaac ggagggagaa 2040 ggattcgaac caacgcaagc acatacatgc tcctaattaa taaaaatata ttaatcccct 2100 taatccagac ttgggtatcc ctccacaagc attatttaat gctaatataa catatataac 2160 aacaaatgtc aatatgtatt tataaggaaa aggatattaa aattattctg agttatataa 2220 ggtagtattc ataatcatcc taaagttgaa gtcgaaaagc ttcaacttta ggaatgagaa 2280 aaaattttta tttgctctaa ttatcatata agtttataaa aagcgcgctc actttatagt 2340 tccccaaaga acacctaaag tattagtaga tttattccat gttcttacag gcaagtaaat 2400 acgttggcca ctgttacctg tataacctac ccaaacatga ccgtcttgtt tcatcacttc 2460 atcataatga attgtttgac ctgcttttaa gactcctgac tgcggcatgc ttctaaatgg 2520 tagtgtgcca cttgttatta tatctgtatt aggtgtgaag ctagctgact ctgatttata 2580 accagtcgtt tatttgtttg ttttccaacc tgtattcggc gttggagtta ctgtaccacc 2640 tgcttttcca tatcctgcgc tctttaagaa aggcattgga tcttgggcag ttgaatttga 2700 aaatgaatta accattettt ggaagtgtaa atgtggtget gtagaataac cagtgettee 2760 agaccaaccg attatttgac cagctttgac ataatctcct actttaacat tatatttact 2820 tagatgcata taccattgtc tatgcactcc atcattttca ataagaccta tttgattacc 2880 tecteegtaa ttaeteeaae eagetteaae tattttteeg ettgaaatag ettttaetgg 2940 tgttccaata ttcataaaaa aatcaactcc gtagtgcata ccgccattta tacctaatgg 3000 ataaggaccg taaccatatc cttttttgta attattcaac cattgtgctg aatgttcatg 3060 tgttgcaget ettaaagetg ttetattttg aaccaggget tttgaagtet etaetteage 3120 tgtattttct actggggctt ttgaagtctc tacttcagct gtattttcta ccggagcttt 3180 tgaagtetet aetteagetg tattttetae tggagetttt gaagteteta etteagetgt 3240 attitictact ggggcttitg aagtototac ticagotgta tittotacog gagottitga 3300 agtetetaet teagetgtat tttetaetgg agettttgaa gtetetaett cagetgtatt 3360 ttctactggg gcttttgaag tctctacttc agctgtattt tctaccggag cttttgaagt 3420 taccggagct tttgaagtct ctacttcagt gtattttcta ctggagcttt tgaagtctct 3480 acttcagctg tattttctac tggagctttt gaagtctcta cttcagctgt attttctact 3540 ggagettttg aagtetetae tteagetgta ttttetaetg gggettttga agtetetaet 3600 tcagctactt tttttgaaac atccatatta cttttttcag aagcatgtgt ttcattttga 3660 atccctccat aaacaataga tgctaaggca aatgtactca gtccaatagc taaaggtctc 3720 gtataataat tgttttttgt tttcttcaac cttaatacct cctaatttat tattacattt 3780 atgagaatag catattttta taaatattac attttttaac gatgaagaaa gtaaaataat 3840 attgattgat tttgatgaaa tatacaaaca cactcaaaaa ttgactttaa gatttttgaa 3900 gaattttcaa aactaaacat tcaagagttc gaagaatttg tgtttcaaaa aatgtctcat 4020

tacacacaat	ctgcttctca	ttttgaatat	agaaataacc	atcagaataa	tgtgcattta	4080
gttggcgtaa	aaaatgaaac	aggtgaagta	ttagctgctt	gtttactgac	tgaggcacgt	4140
tgtttaaagt	tctttaaata	tttctataca	catcgcggtc	cagtcatgaa	ctttaaagac	4200
catgagttag	tcagattttt	ttatgaaaac	ttaacgacct	atctaaaaaa	gcaaaactgc	4260
ttatatgttt	taactgaccc	ttacctgtta	gaaaatattc	gaagttgtga	cggagaaatc	4320
cttgaatctt	atgataacga	aacttttatg	aacgtgatga	atttattagg	ttaccgtcat	4380
caagggttta	ctacaggtta	ttctcaaaca	agtcagatca	gatggttgtc	ggtcttaaac	4440
ctagaaaata	aagatgaaaa	acaattgtta	aaagaaatgg	attatcaaac	acgccgtaat	4500
attaagaaaa	cctatgaaat	gcaggtgaaa	gtccgcgatt	tatcaattaa	tgaaacagat	4560
cgatttttta	aattatttaa	aatggctgaa	gaaaaacatg	gcttcaaata	agttattttg	4620
aaagaatgca	gaaaacatac	gctgataata	gtatgttaaa	gctggcttac	atcgatttag	4680
aagaattatt	agagacacaa	aatgcgaaag	tcgctgagtt	aaatacagat	attgaaaata	4740
ttcaagcggc	attaaaagaa	aaccctaatt	ctaagaaaaa	caaaaataaa	tatgcgcaat	4800
accaaaagca	attagcagca	caagaacgaa	aaattactga	aacgaaaaaa	ttgatagaaa	4860
cagatggacc	tgtattagac	ttagctgcag	cttactatat	ctatacccct	catgaagttt	4920
actacctatc	cagtggttca	aaccctaaat	acaatgccta	tatgggtgcg	tacagactcc	4980
aatgggaaat	gattcaattt	gcgaaaaata	aaggtattaa	tcgctataat	ttttacggta	5040
ttacaggaga	tttcagtgaa	gatgctgaag	atttcggtgt	tcaaaaattc	aaagaaggct	5100
ttaatgccca	tgttgaagaa	tatgtcggcg	acttcattaa	accgattaaa	cctttattt	5160
ataaaattca	tcaattatta	aatagataac	tgaaaattat	ttagtctttg	ttaatcaaat	5220
atgacacctc	aaaatgggtg	tgaagagaac	tatattttca	aaggcgttaa	tctcgacatc	5280
agcgaaggta	aacgttctag	ttttacattc	ttaactacta	agatgctata	atttggttaa	5340
cgaagattat	atgcatatta	agcacctact	tccatcgaaa	atatcgccgg	aagataagac	5400
gactatatta	ttataccatc	tgtaaatata	caagcatata	tacttctgat	aacagaacct	5460
tgtagctgat	gctggctatg	gtagtaaaag	taaggttttg	tttcaaagta	aaaaatatag	5520
ctaaccacta	atttatcatg	tcagtgttca	ctcaacttgc	tagcatgatg	ctaatttcgt	5580
ggcatggcga	aaatccgtag	atctgaagag	atctgcggtt	ctttttatat	agaccgtaaa	5640
tacattcaat	accttttaaa	gtattctttg	ccgtattgat	actttgatac	cttgtctttc	5700
ttactttaat	atgacggtgg	ccttgctcaa	taaggttatt	ccgatatttc	gatgtacaat	5760
gacagtcatg	tttaagttta	aaagctttaa	tgactttagc	catggctacc	ttcgttgaag	5820
gtgcctgatc	tgtaattacc	ttttgaggtt	taccaaattg	tttaatgaga	cgtttgataa	5880
acgcatatgc	tgaatgatta	tctcgttgct	tacgcaagca	aatatctaat	gtatgggttc	5940
	aatactttag	_	-	•		
	aatacttgaa					
	agtatttgct					
	tatgagaaat					
	ggtattctga					
	gattactaaa					
	ctgttaaagt					
	tgtatttcct			attaagatgt	tataccctat	6420
ctttattaat	gctataaacc	gtctgccttg	tgatatc			6457